

Inventors: Huse and Freedman
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*B1
cancel* identify ligands having optimal binding activity. For example, if the collective receptor variant population of this example were screened in the melanophore system, ligand No. 3 would have generated the highest signal since it binds to all seven receptors in the receptor variant population. Ligand No. 7 would give a weaker signal since this ligand binds to three receptors in the receptor variant population. Ligand No. 1 would give a still weaker signal since this ligand binds to two receptors in the receptor variant population. Thus, screening with a collective receptor variant population provides more information about the binding characteristics of the ligand than screening with the parent receptor alone. In addition, ligands that bind weakly to the parent receptor may not have been detectable above background when screened against the parent alone but are detectable when more than one receptor in the receptor variant population binds to the ligand.

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Please delete the table on page 53, lines 1-18, and
substitute therefor:

**Table I. Nucleotide and Amino Acid Sequences of Receptor
Variants of BR96 Antibody**

CDR L1

| | Amino Acid | 26 | 27 | 28 | 29 | 30 | 31 | 32 | 33 |
|------------|------------|-----|------------|------------|------------|------------|------------|-----|-----|
| SEQ ID NO: | | | | | | | | | |
| 1 | Wild type | AGC | TCA | AGT | GTA | AGT | TTC | ATG | AAC |
| 2 | | Ser | Ser | Ser | Val | Ser | Phe | Met | Asn |
| 3 | M131B3-5 | AGC | TCA | AGT | GTA | AGG | TTC | ATG | AAC |
| 4 | | Ser | Ser | Ser | Val | Arg | Phe | Met | Asn |
| 5 | M131B3-6 | AGC | GAG | AGT | GTA | AAT | CTT | ATG | AAC |
| 6 | | Ser | Glu | Ser | Val | Asn | Leu | Met | Asn |
| 7 | M131B3-7 | AGC | TCA | AGT | GTT | AAT | TTC | ATG | AAC |
| 8 | | Ser | Ser | Ser | Val | Asn | Phe | Met | Asn |
| 9 | M131B3-10 | AGC | TCA | ACG | GTA | AGT | TTC | ATG | AAC |
| 10 | | Ser | Ser | Thr | Val | Ser | Phe | Met | Asn |
| 11 | M131B3-11 | AGC | TCA | AGT | GTA | GCG | TAT | ATG | AAC |
| 12 | | Ser | Ser | Ser | Val | Ala | Tyr | Met | Asn |
| 13 | M131B3-12 | AGC | CAG | AGT | GCT | AAG | CAT | ATG | AAC |
| 14 | | Ser | Gln | Ser | Ala | Lys | His | Met | Asn |

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Please delete the table on page 54, lines 1-16, and
substitute therefor:

CDR L2

| | Amino Acid | 49 | 50 | 51 | 52 | 53 | 54 | 55 | 56 |
|------------|------------|-----|-----|------------|------------|------------|-----|-----|-----|
| SEQ ID NO: | | | | | | | | | |
| 15 | Wild type | GCC | ACA | TCC | AAT | TTG | GCT | TCT | GGA |
| 16 | | Ala | Thr | Ser | Asn | Leu | Ala | Ser | Gly |
| 17 | M131B3-5 | GCC | ACA | GAG | AAG | TTG | GCT | TCT | GGA |
| 18 | | Ala | Thr | Glu | Lys | Leu | Ala | Ser | Gly |
| 19 | M131B3-6 | GCC | ACA | GTT | AAT | TTG | GCT | TCT | GGA |
| 20 | | Ala | Thr | Val | Asn | Leu | Ala | Ser | Gly |
| 21 | M131B3-7 | GCC | ACA | GTG | AAT | TTG | GCT | TCT | GGA |
| 22 | | Ala | Thr | Val | Asn | Leu | Ala | Ser | Gly |
| 23 | M131B3-10 | GCC | ACA | TCC | AGG | GCG | GCT | TCT | GGA |
| 24 | | Ala | Thr | Ser | Arg | Ala | Ala | Ser | Gly |
| 25 | M131B3-11 | GCC | ACA | CAG | AAT | TTG | GCT | TCT | GGA |
| 26 | | Ala | Thr | Gln | Asn | Leu | Ala | Ser | Gly |
| 27 | M131B3-12 | GCC | ACA | TCC | AAT | TTG | GCT | TCT | GGA |
| 28 | | Ala | Thr | Ser | Asn | Leu | Ala | Ser | Gly |